RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/552,896Source: 1FwpDate Processed by STIC: 9/01/2006

ENTERED

CRF Errors Edited by the STIC Systems Branch

Serial	Number: 10/552, 896	CRF Edit Date: 9/01/2006 Edited by: DA
	Realigned nucleic acid/amino acid numbers/text in cases where the sequence text "wrapped" to the next line	
	Corrected the SEQ ID NO. Sequence numbers edited were:	
	Inserted or corrected a nucleic number at the e NO's edited:	end of a nucleic line. SEQ ID
_/	Deleted: invalid beginning/end-of-file text	; page numbers
	Inserted mandatory headings/numeric identifie	ers, specifically:
	Moved responses to same line as heading/nume	eric identifier, specifically:
	Other:	

Revised 09/09/2003



- 45 40

IFWP

RAW SEQUENCE LISTING DATE: 09/01/2006 PATENT APPLICATION: US/10/552,896 TIME: 12:13:41

Input Set : A:\pto.da.txt

3 <110> APPLICANT: Neose Technologies, Inc.

```
4
             DeFrees, Shawn
             Zopf, David
     5
             Bayer, Robert
             Hakes, David
     8
             Chen, Xi
             Bowe, Caryne
     9
    11 <120> TITLE OF INVENTION: GLYCOPEGYLATION METHODS AND PROTEINS/PEPTIDES PRODUCED BY
THE
    12
             METHODS
    14 <130> FILE REFERENCE: 040853-5051-US01
    16 <140> CURRENT APPLICATION NUMBER: US 10/552,896
    17 <141> CURRENT FILING DATE: 2005-10-11
    19 <150> PRIOR APPLICATION NUMBER: US 10/411,012
    20 <151> PRIOR FILING DATE: 2003-04-09
    22 <150> PRIOR APPLICATION NUMBER: US 10/411,026
    23 <151> PRIOR FILING DATE: 2003-04-09
    25 <150> PRIOR APPLICATION NUMBER: US 10/410,962
    26 <151> PRIOR FILING DATE: 2003-04-09
    28 <150> PRIOR APPLICATION NUMBER: US 10/411,049
    29 <151> PRIOR FILING DATE: 2003-04-09
    31 <150> PRIOR APPLICATION NUMBER: US 10/410,930
    32 <151> PRIOR FILING DATE: 2003-04-09
    34 <150> PRIOR APPLICATION NUMBER: US 10/410,897
    35 <151> PRIOR FILING DATE: 2003-04-09
    37 <150> PRIOR APPLICATION NUMBER: US 10/410,997
    38 <151> PRIOR FILING DATE: 2003-04-09
    40 <150> PRIOR APPLICATION NUMBER: US 10/411,044
    41 <151> PRIOR FILING DATE: 2003-04-09
    43 <150> PRIOR APPLICATION NUMBER: US 10/410,980
    44 <151> PRIOR FILING DATE: 2003-04-09
    46 <150> PRIOR APPLICATION NUMBER: US 10/410,945
    47 <151> PRIOR FILING DATE: 2003-04-09
    49 <150> PRIOR APPLICATION NUMBER: US 10/410,913
    50 <151> PRIOR FILING DATE: 2003-04-09
    52 <150> PRIOR APPLICATION NUMBER: US 10/411,037
    53 <151> PRIOR FILING DATE: 2003-04-09
    55 <150> PRIOR APPLICATION NUMBER: US 10/411,043
    56 <151> PRIOR FILING DATE: 2003-04-09
    58 <150> PRIOR APPLICATION NUMBER: PCT US2004/011494
    59 <151> PRIOR FILING DATE: 2004-04-09
    61 <160> NUMBER OF SEQ ID NOS: 75
    63 <170> SOFTWARE: PatentIn version 3.2
    65 <210> SEQ ID NO: 1
```

Input Set : A:\pto.da.txt

Output Set: N:\CRF4\09012006\J552896.raw

```
66 <211> LENGTH: 525
67 <212> TYPE: DNA
68 <213> ORGANISM: Homo sapiens
70 <400> SEQUENCE: 1
71 accccctgg gcctgccag ctccctgccc cagagettcc tgctcaagtg cttagagcaa
                                                                         60
73 gtgaggaaga tccagggcga tggcgcagcg ctccaggaga agctgtgtgc cacctacaag
                                                                        120
75 ctgtgccacc ccgaggaget ggtgctgctc ggacactetc tgggcatecc ctgggctccc
                                                                        180
77 ctgagcaget geeceageea ggeeetgeag etggeagget gettgageea acteeatage
                                                                        240
79 gqccttttcc tctaccaggg gctcctgcag gccctggaag ggatctcccc cgagttgggt
                                                                        300
81 cccaccttgg acacactgca gctggacgtc gccgactttg ccaccaccat ctggcagcag
                                                                        360
83 atggaagaac tgggaatggc ccctgccctg cagcccaccc agggtgccat gccggccttc
                                                                        420
85 gcctctgctt tccagcgccg ggcaggaggg gtcctggttg cctcccatct gcagagcttc
                                                                        480
87 ctggaggtgt cgtaccgcgt tctacgccac cttgcccagc cctga
                                                                        525
89 <210> SEQ ID NO: 2
90 <211> LENGTH: 174
91 <212> TYPE: PRT
92 <213> ORGANISM: Homo sapiens
94 <400> SEQUENCE: 2
95 Thr Pro Leu Gly Pro Ala Ser Ser Leu Pro Gln Ser Phe Leu Leu Lys
98 Cys Leu Glu Gln Val Arg Lys Ile Gln Gly Asp Gly Ala Ala Leu Gln
               20
                                   25
101 Glu Lys Leu Cys Ala Thr Tyr Lys Leu Cys His Pro Glu Glu Leu Val
104 Leu Leu Gly His Ser Leu Gly Ile Pro Trp Ala Pro Leu Ser Ser Cys
                            55
107 Pro Ser Gln Ala Leu Gln Leu Ala Gly Cys Leu Ser Gln Leu His Ser
110 Gly Leu Phe Leu Tyr Gln Gly Leu Leu Gln Ala Leu Glu Gly Ile Ser
                    85
                                        90
113 Pro Glu Leu Gly Pro Thr Leu Asp Thr Leu Gln Leu Asp Val Ala Asp
116 Phe Ala Thr Thr Ile Trp Gln Gln Met Glu Glu Leu Gly Met Ala Pro
                                                     125
117
            115
                                120
119 Ala Leu Gln Pro Thr Gln Gly Ala Met Pro Ala Phe Ala Ser Ala Phe
                            135
122 Gln Arg Arg Ala Gly Gly Val Leu Val Ala Ser His Leu Gln Ser Phe
                        150
                                            155
123 145
125 Leu Glu Val Ser Tyr Arg Val Leu Arg His Leu Ala Gln Pro
128 <210> SEQ ID NO: 3
129 <211> LENGTH: 1733
130 <212> TYPE: DNA
131 <213> ORGANISM: Homo sapiens
133 <400> SEQUENCE: 3
134 gegeetetta tgtacceaca aaaatetatt ttcaaaaaag ttgetetaag aatatagtta
                                                                          60
```

136 tcaaqttaag taaaatgtca atagcctttt aatttaattt ttaattgttt tatcattctt

138 tgcaataata aaacattaac tttatacttt ttaatttaat gtatagaata gagatataca

140 taggatatgt aaatagatac acagtgtata tgtgattaaa atataatggg agattcaatc

120

180

240

Input Set : A:\pto.da.txt

```
142 agaaaaaagt ttctaaaaag gctctggggt aaaagaggaa ggaaacaata atgaaaaaaa
 144 tqtqqtqaqa aaaacagctq aaaacccatq taaaqagtqt ataaagaaaq caaaaagaga
                                                                          360
 146 aqtaqaaaqt aacacagggg catttggaaa atgtaaacga gtatgttccc tatttaaggc
                                                                          420
 148 taggcacaaa gcaaggtctt cagagaacct ggagcctaag gtttaggctc acccatttca
                                                                          480
 150 accagtctag cagcatctgc aacatctaca atggccttga cctttgcttt actggtggcc
                                                                          540
 152 ctcctggtgc tcagctgcaa gtcaagctgc tctgtgggct gtgatctgcc tcaaacccac
                                                                          600
 154 agectgggta geaggaggae ettgatgete etggeaeaga tgaggagaat etetettte
                                                                          660
                                                                          720
 156 tectgettga aggacagaca tgactttgga tttccccagg aggagtttgg caaccagtte
 158 caaaaggetg aaaccatece tgteeteeat gagatgatee ageagatett caatetette
                                                                          780
 160 agcacaaagg actcatctgc tgcttgggat gagaccctcc tagacaaatt ctacactgaa
                                                                          840
 162 ctctaccage agctgaatga cctggaagcc tgtgtgatac agggggtggg ggtgacagag
                                                                          900
 164 actcccctga tgaaggagga ctccattctg gctgtgagga aatacttcca aagaatcact
                                                                          960
 166 ctctatctga aagagaagaa atacagccct tgtgcctggg aggttgtcag agcagaaatc
                                                                         1020
 168 atgagatett tttetttgte aacaaacttg caagaaagtt taagaagtaa ggaatgaaaa
 170 ctggttcaac atggaaatga ttttcattga ttcgtatgcc agctcacctt tttatgatct
                                                                         1140
                                                                         1200
 172 gccatttcaa agactcatgt ttctgctatg accatgacac gatttaaatc ttttcaaatg
 174 tttttaggag tattaatcaa cattgtattc agctcttaag gcactagtcc cttacagagg
                                                                         1260
 176 accatgctga ctgatccatt atctatttaa atatttttaa aatattattt atttaactat
. 178-ttataaaaga acttattttt gttcatatta tgtcatgtgc acctttgcac agtggttaat
                                                                         1380
 180 graataaaat gigitetiig tattiggtaa attiattiig igitgiteat igaaettiig
                                                                         1440
 182 ctatggaact tttgtacttg tttattcttt aaaatgaaat tccaagccta attgtgcaac
 184 ctgattacag aataactggt acacttcatt tgtccatcaa tattatattc aagatataag
                                                                         1560
 186 taaaaataaa ctttctgtaa accaagttgt atgttgtact caagataaca gggtgaacct
                                                                         1620
 188 aacaaataca attctqctct cttqtqtatt tqatttttqt atqaaaaaaa ctaaaaatgg
                                                                        1680
                                                                         1733
 190 taatcatact taattatcag ttatggtaaa tggtatgaag agaagaagga acg
 192 <210> SEQ ID NO: 4
 193 <211> LENGTH: 188
 194 <212> TYPE: PRT
 195 <213> ORGANISM: Homo sapiens
 197 <400> SEQUENCE: 4
 198 Met Ala Leu Thr Phe Ala Leu Leu Val Ala Leu Leu Val Leu Ser Cys
 199 1
 201 Lys Ser Ser Cys Ser Val Gly Cys Asp Leu Pro Gln Thr His Ser Leu
 202
                                      25
                 20
 204 Gly Ser Arg Arg Thr Leu Met Leu Leu Ala Gln Met Arg Arg Ile Ser
 205
 207 Leu Phe Ser Cys Leu Lys Asp Arg His Asp Phe Gly Phe Pro Gln Glu
 208
                              55
 210 Glu Phe Gly Asn Gln Phe Gln Lys Ala Glu Thr Ile Pro Val Leu His
 213 Glu Met Ile Gln Gln Ile Phe Asn Leu Phe Ser Thr Lys Asp Ser Ser
                                          90
 216 Ala Ala Trp Asp Glu Thr Leu Leu Asp Lys Phe Tyr Thr Glu Leu Tyr
                                      105
                 100
 219 Gln Gln Leu Asn Asp Leu Glu Ala Cys Val Ile Gln Gly Val Gly Val
                                  120
 222 Thr Glu Thr Pro Leu Met Lys Glu Asp Ser Ile Leu Ala Val Arg Lys
                              135
 225 Tyr Phe Gln Arg Ile Thr Leu Tyr Leu Lys Glu Lys Lys Tyr Ser Pro
```

Input Set : A:\pto.da.txt

```
226 145
                         150
                                             155
 228 Cys Ala Trp Glu Val Val Arg Ala Glu Ile Met Arg Ser Phe Ser Leu
                                         170
                     165
 231 Ser Thr Asn Leu Gln Glu Ser Leu Arg Ser Lys Glu
 232
                 180
 234 <210> SEQ ID NO: 5
 235 <211> LENGTH: 757
 236 <212> TYPE: DNA
 237 <213> ORGANISM: Homo sapiens
 239 <400> SEQUENCE: 5
 240 atgaccaaca agtgtctcct ccaaattgct ctcctgttgt gcttctccac tacagctctt
 242 tccatgagct acaacttgct tggattccta caaagaagca gcaattttca gtgtcagaag
                                                                        120
 244 ctcctgtggc aattgaatgg gaggcttgaa tattgcctca aggacaggat gaactttgac
                                                                        180
 246 atccctgagg agattaagca gctgcagcag ttccagaagg aggacgccgc attgaccatc
                                                                        240
 248 tatgagatgc tccagaacat ctttgctatt ttcagacaag attcatctag cactggctgg
                                                                        300
                                                                        360
 250 aatgagacta ttgttgagaa cctcctggct aatgtctatc atcagataaa ccatctgaag
 252 acagtcctgg aagaaaaact ggagaaagaa gattttacca ggggaaaact catgagcagt
                                                                        420
 254 ctgcacctga aaagatatta tgggaggatt ctgcattacc tgaaggccaa ggagtacagt
                                                                        480
.. 256 cactytweet ggaedatagt dagagtggaa atdetaagga acttttactt dabtaadaga ...
                                                                        600 - Caraman - Cara - Caraman
- 258 ottacaggit accidegaaa eigaagatei eelageelgi ceeleiggga eiggacaalt
 260 gcttcaagca ttcttcaacc agcagatgct gtttaagtga ctgatggcta atgtactgca
 720
                                                                        757
 264 ttaaatttta ttttggaaaa taaattattt ttggtgc
 266 <210> SEQ ID NO: 6
 267 <211> LENGTH: 187
  268 <212> TYPE: PRT
  269 <213> ORGANISM: Homo sapiens
  271 <400> SEQUENCE: 6
 272 Met Thr Asn Lys Cys Leu Leu Gln Ile Ala Leu Leu Cys Phe Ser
 273 1
 275 Thr Thr Ala Leu Ser Met Ser Tyr Asn Leu Leu Gly Phe Leu Gln Arg
 276
 278 Ser Ser Asn Phe Gln Cys Gln Lys Leu Leu Trp Gln Leu Asn Gly Arg
 279
  281 Leu Glu Tyr Cys Leu Lys Asp Arg Met Asn Phe Asp Ile Pro Glu Glu
  282
                             55
  284 Ile Lys Gln Leu Gln Gln Phe Gln Lys Glu Asp Ala Ala Leu Thr Ile
                         70
  287 Tyr Glu Met Leu Gln Asn Ile Phe Ala Ile Phe Arg Gln Asp Ser Ser
                     85
  290 Ser Thr Gly Trp Asn Glu Thr Ile Val Glu Asn Leu Leu Ala Asn Val
  291
                 100
                                     105
  293 Tyr His Gln Ile Asn His Leu Lys Thr Val Leu Glu Glu Lys Leu Glu
             115
                                 120
                                                     125
  296 Lys Glu Asp Phe Thr Arg Gly Lys Leu Met Ser Ser Leu His Leu Lys
                             135
  299 Arg Tyr Tyr Gly Arg Ile Leu His Tyr Leu Lys Ala Lys Glu Tyr Ser
  302 His Cys Ala Trp Thr Ile Val Arg Val Glu Ile Leu Arg Asn Phe Tyr
```

Input Set : A:\pto.da.txt

```
303
                     165
                                         170
                                                              175
305 Phe Ile Asn Arg Leu Thr Gly Tyr Leu Arg Asn
                 180
308 <210> SEQ ID NO: 7
309 <211> LENGTH: 1332
310 <212> TYPE: DNA
311 <213> ORGANISM: Homo sapiens
313 <400> SEQUENCE: 7
                             5
314 atggtetece aggeeeteag geteetetge ettetgettg ggetteaggg etgeetgget
                                                                           60
316 gcagtetteg taacecagga ggaageecae ggegteetge aeeggegeeg gegegeeaae
                                                                          120
318 gcgttcctgg aggagctgcg gccgggctcc ctggagaggg agtgcaagga ggagcagtgc
                                                                          180
                                                                          240
320 teettegagg aggeeeggga gatetteaag gaegeggaga ggaegaaget gttetggatt
322 tettacagtg atggggacca gtgtgeetca agtecatgee agaatggggg eteetgeaag
                                                                          300
324 gaccagetee agteetatat etgettetge etceetgeet tegagggeeg gaactgtgag
                                                                          360
                                                                          420
326 acgcacaagg atgaccagct gatctgtgtg aacgagaacg gcggctgtga gcagtactgc
                                                                          480
328 agtgaccaca cgggcaccaa gcgctcctgt cggtgccacg aggggtactc tctgctggca
                                                                          540
330 gacggggtgt cctgcacacc cacagttgaa tatccatgtg gaaaaatacc tattctagaa
332 aaaagaaatg ccagcaaacc ccaaggccga attgtggggg gcaaggtgtg ccccaaaggg
                                                                          600
.334 gagtgtgcat, ggcaggtcot gttgttgdtg aatggagctc agttgtgtgg ggggaccetg
                                                                         ·660
                                                                          720
336 ateaacacca tetgggtggt etecgeggee-eactgttteg acaaaateaa gaactggagg
                                                                          780
338 aacctgatcg cggtgctggg cgagcacgac ctcagcgagc acgacgggga tgagcagagc
                                                                          840
340 eggegggtgg egeaggteat cateceeage aegtaegtee egggeaceae caaceaegae
                                                                          900
342 ategegetge teegeetgea eeageeegtg gteeteactg accatgtggt geeectetge
344 etgecegaac ggacgttete tgagaggacg etggeetteg tgegettete attggteage
                                                                          960
346 ggctggggcc agctgctgga ccgtggcgcc acggccctgg agctcatggt gctcaacgtg
                                                                         1020
348 ccccggctga tgacccagga ctgcctgcag cagtcacgga aggtgggaga ctccccaaat
                                                                         1080
350 atcacggagt acatgttctg tgccggctac tcggatggca gcaaggactc ctgcaagggg
                                                                         1140
352 gacagtggag gcccacatgc cacccactac cggggcacgt ggtacctgac gggcatcgtc
                                                                         1200
354 agetggggce agggetgege aacegtggge caetttgggg tgtacaccag ggteteceag
                                                                         1260
356 tacatcgagt ggctgcaaaa gctcatgcgc tcagagccac gcccaggagt cctcctgcga
                                                                         1320
                                                                         1332
358 gccccatttc cc
 360 <210> SEQ ID NO: 8
 361 <211> LENGTH: 444
 362 <212> TYPE: PRT
 363 <213> ORGANISM: Homo sapiens
 365 <400> SEQUENCE: 8
 366 Met Val Ser Gln Ala Leu Arg Leu Leu Cys Leu Leu Leu Gly Leu Gln
367 1
                                         10
 368 Gly Cys Leu Ala Ala Val Phe Val Thr Gln Glu Glu Ala His Gly Val
371 Leu His Arg Arg Arg Ala Asn Ala Phe Leu Glu Glu Leu Arg Pro
372
                                 40
374 Gly Ser Leu Glu Arg Glu Cys Lys Glu Glu Gln Cys Ser Phe Glu Glu
375
        50
                             55
377 Ala Arg Glu Ile Phe Lys Asp Ala Glu Arg Thr Lys Leu Phe Trp Ile
378 65
                         70
                                             75
380 Ser Tyr Ser Asp Gly Asp Gln Cys Ala Ser Ser Pro Cys Gln Asn Gly
383 Gly Ser Cys Lys Asp Gln Leu Gln Ser Tyr Ile Cys Phe Cys Leu Pro
```

VERIFICATION SUMMARY

DATE: 09/01/2006

PATENT APPLICATION: US/10/552,896

TIME: 12:13:42

Input Set : A:\pto.da.txt

Output Set: N:\CRF4\09012006\J552896.raw

Services

Raw Sequence Listing before editing, for reference only



IFWP

RAW SEQUENCE LISTING DATE: 08/28/2006 PATENT APPLICATION: US/10/552,896 TIME: 10:22:21

Input Set : A:\040853-01-5051US01 seq list.TXT

Output Set: N:\CRF4\08282006\J552896.raw

```
3 <110 > APPLICANT: Neose Technologies, Inc.
       DeFrees, Shawn
         Zopf, David
         Bayer, Robert
 6
         Hakes, David
 7
 8
         Chen, Xi
         Bowe, Caryne
11 <120> TITLE OF INVENTION: GLYCOPEGYLATION METHODS AND PROTEINS/PEPTIDES PRODUCED BY
         METHODS
14 <130> FILE REFERENCE: 040853-5051-US01
                                                           Does Not Comply
Corrected Diskette Needed
16 <140> CURRENT APPLICATION NUMBER: US 10/552,896
17 <141> CURRENT FILING DATE: 2005-10-11
19 <150> PRIOR APPLICATION NUMBER: US 10/411,012
20 <151> PRIOR FILING DATE: 2003-04-09
22 <150> PRIOR APPLICATION NUMBER: US 10/411,026
23 <151> PRIOR FILING DATE: 2003-04-09
25 <150> PRIOR APPLICATION NUMBER: US 10/410,962
26 <151> PRIOR FILING DATE: 2003-04-09
28 <150> PRIOR APPLICATION NUMBER: US 10/411,049
29 <151> PRIOR FILING DATE: 2003-04-09
31 <150> PRIOR APPLICATION NUMBER: US 10/410,930
32 <151> PRIOR FILING DATE: 2003-04-09
34 <150> PRIOR APPLICATION NUMBER: US 10/410,897
35 <151> PRIOR FILING DATE: 2003-04-09
37 <150> PRIOR APPLICATION NUMBER: US 10/410,997
38 <151> PRIOR FILING DATE: 2003-04-09
40 <150> PRIOR APPLICATION NUMBER: US 10/411,044
41 <151> PRIOR FILING DATE: 2003-04-09
43 <150> PRIOR APPLICATION NUMBER: US 10/410,980
44 <151> PRIOR FILING DATE: 2003-04-09
46 <150> PRIOR APPLICATION NUMBER: US 10/410,945
47 <151> PRIOR FILING DATE: 2003-04-09
49 <150> PRIOR APPLICATION NUMBER: US 10/410,913
50 <151> PRIOR FILING DATE: 2003-04-09
52 <150> PRIOR APPLICATION NUMBER: US 10/411,037
53 <151> PRIOR FILING DATE: 2003-04-09
55 <150> PRIOR APPLICATION NUMBER: US 10/411,043
56 <151> PRIOR FILING DATE: 2003-04-09
58 <150> PRIOR APPLICATION NUMBER: PCT US2004/011494
59 <151> PRIOR FILING DATE: 2004-04-09
61 <160> NUMBER OF SEQ ID NOS: 75
63 <170> SOFTWARE: PatentIn version 3.2
```

THE

Input Set : A:\040853-01-5051US01 seq list.TXT

Output Set: N:\CRF4\08282006\J552896.raw

ERRORED SEQUENCES

```
4804 <210> SEQ ID NO: 75
     4805 <211> LENGTH: 195
     4806 <212> TYPE: PRT
     4807 <213> ORGANISM: Homo sapiens
     4809 <400> SEQUENCE: 75
    4810 Met Ala Leu Leu Phe Pro Leu Leu Ala Ala Leu Val Met Thr Ser Tyr
     4811 1
     4813 Ser Pro Val Gly Ser Leu Gly Cys Asp Leu Pro Gln Asn His Gly Leu
     4816 Leu Ser Arg Asn Thr Leu Val Leu Leu His Gln Met Arg Arg Ile Ser
     4817
     4819 Pro Phe Leu Cys Leu Lys Asp Arg Arg Asp Phe Arg Phe Pro Gln Glu
     4822 Met Val Lys Gly Ser Gln Leu Gln Lys Ala His Val Met Ser Val Leu
     4823 65
                             70
                                                 75
     4825 His Glu Met Leu Gln Gln Ile Phe Ser Leu Phe His Thr Glu Arg Ser
     4828 Ser Ala Ala Trp Asn Met Thr Leu Leu Asp Gln Leu His Thr Gly Leu
     4829
                     100
     4831 His Gln Gln Leu Gln His Leu Glu Thr Cys Leu Leu Gln Val Val Gly
     4832
                 115
                                     120
     4834 Glu Gly Glu Ser Ala Gly Ala Ile Ser Ser Pro Ala Leu Thr Leu Arg
     4835
                                 135
                                                     140
     4837 Arg Tyr Phe Gln Gly Ile Arg Val Tyr Leu Lys Glu Lys Lys Tyr Ser
     4838 145
                             150
                                                 155
     4840 Asp Cys Ala Trp Glu Val Val Arg Met Glu Ile Met Lys Ser Leu Phe
                         165
                                             170
    4843 Leu Ser Thr Asn Met Gln Glu Arg Leu Arg Ser Lys Asp Arg Asp Leu
    4844
                     180
                                                             190
    4846 Gly Ser Ser
    4847
    4862 (1996240.1
E--> 4863 (continued).(continued).(continued).(continued).(continued).(continued).
(continued).(continued).(continued).(continued).(continued).(continued).
(continued).(continued).(conti
E--> 4864 /htinued).(continued).(continued).(continued).(continued).
E--> 4866 1
E--> 4869 1
                                        deleted
```

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 08/28/2006 PATENT APPLICATION: US/10/552,896 TIME: 10:22:22

Input Set : A:\040853-01-5051US01 seq list.TXT

Output Set: N:\CRF4\08282006\J552896.raw

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:75; Line(s) 4863

VERIFICATION SUMMARY DATE: 08/28/2006
PATENT APPLICATION: US/10/552,896 TIME: 10:22:22

Input Set : A:\040853-01-5051US01 seq list.TXT
Output Set: N:\CRF4\08282006\J552896.raw

L:4863 M:360 E: Sequence data overflow, line data truncated, for SEQ ID#:75

L:4863 M:333 E: Wrong sequence grouping, Amino acids not in groups!

L:4863 M:330 E: (2) Invalid Amino Acid Designator, NUMBER OF INVALID KEYS:1

L:4864 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:75

L:4864 M:333 E: Wrong sequence grouping, Amino acids not in groups!

L:4864 M:330 E: (2) Invalid Amino Acid Designator, NUMBER OF INVALID KEYS:1

M:332 Repeated in SeqNo=75

1. 1/2 1. 1/4

L:4869 M:252 E: No. of Seq. differs, <211> LENGTH:Input:195 Found:197 SEQ:75